

**General Stats** 

Cutadapt

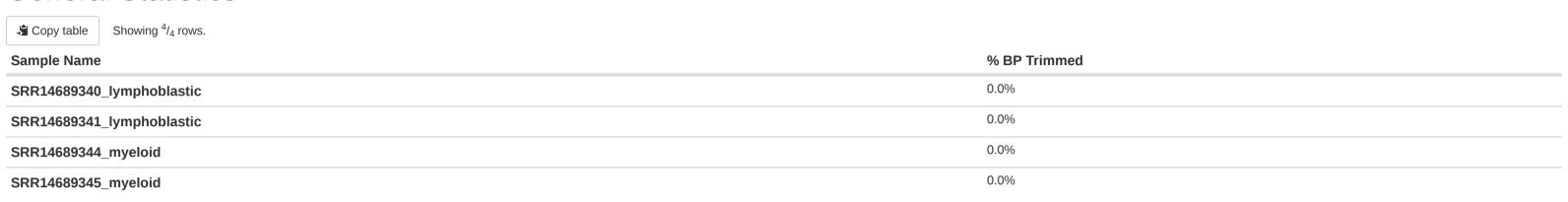


A modular tool to aggregate results from bioinformatics analyses across many samples into a single report.

Report generated on 2022-04-23, 15:56 based on data in: /home/khoidnyds/RNAseq\_old/3.cutadapt

Watch a tutorial video (6:06)

## **General Statistics**



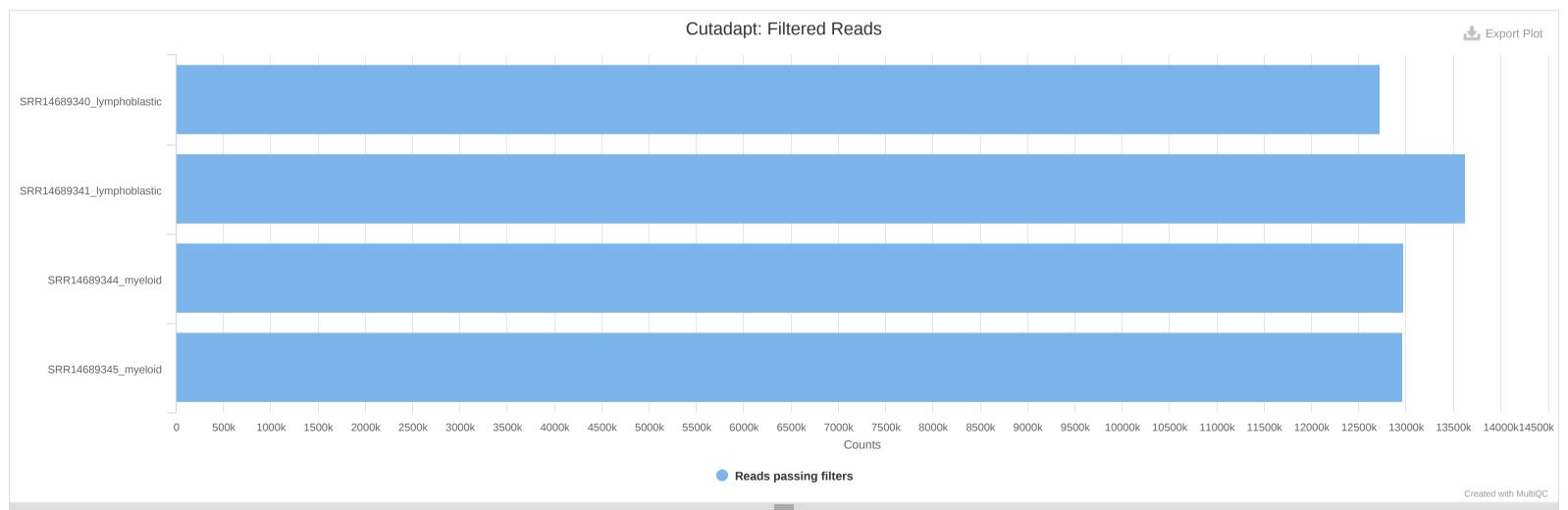
## Cutadapt

Cutadapt is a tool to find and remove adapter sequences, primers, poly-A tails and other types of unwanted sequence from your high-throughput sequencing reads. DOI: 10.14806/ej.17.1.200.

## Filtered Reads

This plot shows the number of reads (SE) / pairs (PE) removed by Cutadapt.







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